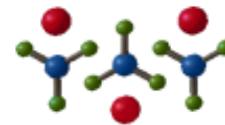




Stony Brook University



Oganov's

COMPUTATIONAL MATERIALS DISCOVERY LABORATORY

# Tutorial 8: Introduction to USPEX

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# Useful links:

- References, where the method was exhaustively described: <http://uspex.stonybrook.edu/uspex.html>
- USPEX manual:
  - HTML: [http://han.ess.sunysb.edu/uspex\\_manual/](http://han.ess.sunysb.edu/uspex_manual/)
  - PDF: [http://han.ess.sunysb.edu/uspex\\_manual/uspex\\_manual.pdf](http://han.ess.sunysb.edu/uspex_manual/uspex_manual.pdf)
- USPEX utilities: <http://han.ess.sunysb.edu/>

# How to start your calculations

1. Prepare INPUT.txt file.
2. Prepare files in **Specific/** directory (for VASP: INCAR\_1,...,INCAR\_N, POSCAR\_A, POSCAR\_B, etc.)
3. If necessary, prepare additional files, which are described in the manual.
4. Start calculation: **USPEX -r**

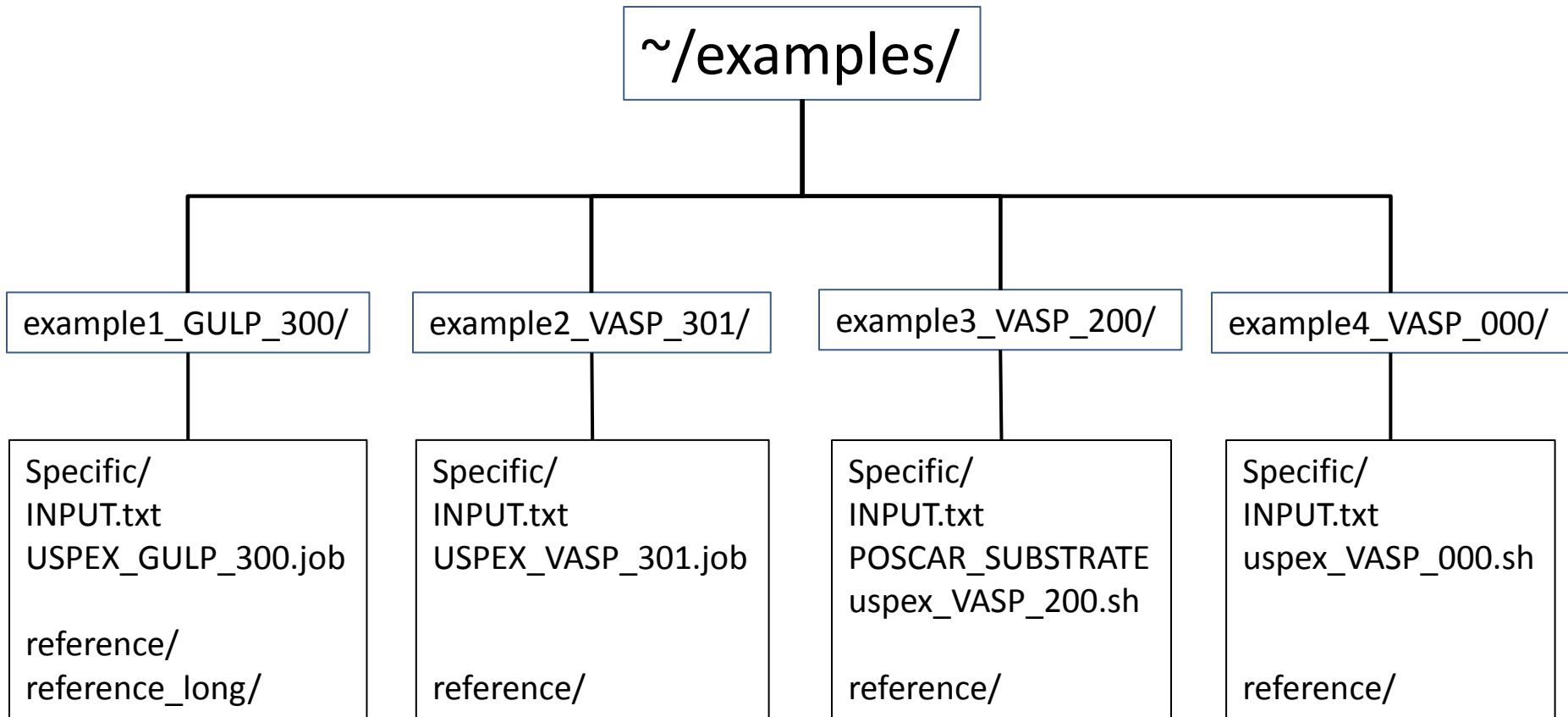
# Connection information:

- `ssh USPEXBNL@qsh.ess.sunysb.edu`  
Password: \*\*\*\*\* (please enter the password correctly, otherwise we will be blocked)
- `cd ~/workshop`
- `mkdir <your userid on nano.bnl.gov>`
- `cd <your userid on nano.bnl.gov>`
- `USPEX -h`

# Examples for today:

- Bulk  $\text{Mg}_4\text{Al}_8\text{O}_{16}$  (fixed composition, calculationType=300, GULP)
- Bulk Si-C (variable composition, calculationType=301, VASP)
- $\text{O}_2$  @ Si (100) (fixed composition, calculationType=200, VASP)
- $\text{C}_{13}$  nanoparticles (fixed composition, calculationType=000, VASP)

# Examples file system



# Examples for today:

- Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (fixed composition, calculationType=300, GULP)
- Bulk Si-C (variable composition, calculationType=301, VASP)
- O<sub>2</sub> @ Si (100) (fixed composition, calculationType=200, VASP)
- C<sub>13</sub> nanoparticles (fixed composition, calculationType=000, VASP)

# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

MgAl<sub>2</sub>O<sub>4</sub> (28 atoms/cell) at 100 GPa pressure.  
Variable-cell calculation using Buckingham  
potentials, GULP code. This test has direct bearing  
on the physics of the Earth's interior!

# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## INPUT.txt:

```
1 ****  
2 *      TYPE OF RUN AND SYSTEM          *  
3 ****  
4 USPEX   : calculationMethod (USPEX, VCNEB, META)  
5 300    : calculationType (dimension: 0-3; molecule: 0/1;  
6 1      : optType (1=enthalpy, 2=volume, 3=hardness, 4=st  
7  
8 % symmetries  
9 2-230  
10 % endSymmetries  
11  
12 % atomType  
13 Mg Al O  
14 % EndAtomType  
15  
16 % numSpecies  
17 4 8 16  
18 % EndNumSpecies  
19  
20 % valences  
21 2 3 2  
22 % endValences  
23
```

### > variable optType

*Meaning:* This variable allows you to specify the quantity to be optimized.

Possible values (characters):

- enthalpy (or "1") — to find the stable phases
- volume (or "2") — volume minimization (to find the most compact structure)
- hardness (or "3") — hardness maximization (to find the hardest structure)
- struc\_order (or "4") — maximization of the degree of order in the structure
- aver\_dist (or "5") — maximization of average distance between atoms
- diel\_sus (or "6") — maximization of the static dielectric susceptibility (only for VASP and GULP)
- gap (or "7") — maximization of the band gap (only for VASP)
- diel\_gap (or "8") — maximization of electrical energy storage capacity (only for VASP)
- mag\_moment (or "9") — maximization of the magnetization (only for VASP)
- quasientropy (or "10") — maximization of structural quasientropy

### > variable calculationType

*Meaning:* Specifies type of calculation, i.e., whether the structure of a bulk crystal, nanoparticle, or surface is to be predicted. This variable consists of three indices: *dimensionality*, *molecularity*, and *compositional variability*:

- dimensionality:

- "3" — bulk crystals
- "2" — surfaces, "-2" — 2D-crystals
- "1" — polymers
- "0" — nanoparticles

- molecularity:

- "0" — non-molecular
- "1" — molecular calculations

- variability of chemical composition in the calculation:

- "0" — fixed composition
- "1" — variable composition

*Default:* 300

*Format:*

```
301 : calculationType
```

*Note:* If *calculationType*=310, i.e., a prediction for a molecular crystal is to be performed, then USPEX expects you to provide files MOL\_1, MOL\_2, ... with molecular geometries for all types of molecules, and these molecules will be placed in the newly generated structures as whole objects. Available options: 300, 301, 310, 000, 200, 201, -200 (and not yet released: 110, 311.)



# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## INPUT.txt:

Possible space groups for crystals

```

1 ****
2 *      TYPE OF RUN AND SYSTEM      *
3 ****
4 USPEX  : calculationMethod (USPEX, VCNEB, META)
5 300   : calculationType (dimension: 0-3; molecule: 0/1; varcomp: 0/1)
6 1     : optType (1=enthalpy, 2=volume, 3=hardness, 4=struc_order, 5=aver_dist)
7
8 % symmetries
9 2-230
10 % endSymmetries
11
12 % atomType
13 Mg Al O
14 % EndAtomType
15
16 % numSpecies
17 4 8 16
18 % EndNumSpecies
19
20 % valences
21 2 3 2
22 % endValences

```

▷ variable `atomType`

*Meaning:* Describes the identities of each type of atom.

*Default:* none, must specify explicitly

*Format:*

If you prefer to use the atomic numbers from Mendeleev's Periodic Table of the Elements, specify:

```
% atomType
12 14 8
% EndAtomType
```

Or, if you prefer to use atomic names, specify:

```
% atomType
Mg Si O
% EndAtomType
```

You can alternatively specify the full names of the elements, for example:

```
% atomType
Magnesium Silicon Oxygen
% EndAtomType
```

Number of atoms of each type

Valences of each type of atom

### 10 Appendix: List of 230 space groups

1	P1	2	P-1	3	P2	4	P21	5	C2
6	Pm	7	P6	8	Cm	14	C2/m	9	P2/m
11	P21/m	12	C2/m	13	P21/c	15	C2/c	10	P21/c
16	P222	17	P2221	18	P2221	19	P222121	20	C2221
21	C222	22	F222	23	I222	24	I222121	25	Pmm2
26	Pmc21	27	Pcc2	28	Pma2	29	Pca21	30	Pnc2
31	Pmn21	32	Pba2	33	Pmd21	34	Pm2	35	Cnn2
36	Cmcm2	37	Pcd2	38	Pcm2	39	Acm2	40	Cnn2
41	Aba2	42	Fmm2	43	Fdd2	44	Innm2	45	Bsc2
46	Imm2	47	Pmmm	48	Pmmn	49	Pcmn	50	Pbsm
51	Pmma	52	Pmma	53	Pmma	54	Peca	55	Pbam
56	Pcca	57	Pbcm	58	Pcmn	59	Pnma	60	Pbhn
61	Pbca	62	Pbcm	63	Cmcm	64	Cmca	65	Cmmn
66	Cmcm	67	Cmcm	68	Cmca	69	H2d	70	H2hd
71	Immm	72	Ibam	73	Ibec	74	Imma	75	P4
76	P41	77	P43	78	P43	79	I4	80	I41
81	P4	82	I-4	83	P4/m	84	P42/m	85	P4/n
86	P42/m	87	I-4/m	88	I4/m	89	P422	90	P422
91	P422	92	I422	93	I422	94	I422	95	P422
96	P422	97	I422	98	I422	99	P4mm	100	Pbhm
101	P4cm	102	P4gm	103	P4c	104	P4nc	105	P4gmc
106	P4bc	107	I4m	108	I4cm	109	I4md	110	I4red
111	P-42m	112	P-43m	113	P-42_m	114	P-42_m2	115	P-42_m2

# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## INPUT.txt:

```
24 ****
25 *          POPULATION *
26 ****
27 5   : populationSize (how many individuals per generation)
28 5   : initialPopSize (how many individuals in the first generation)
29 3   : numGenerations (how many generations shall be calculated)
30 3   : stopCrit (max number of generations with the same best structure)
31 0   : reoptOld (should the old structures be reoptimized)
32 0.6 : bestFrac (What fraction of current generation shall be used to produce the next generation)
33 ****
34 *          VARIATION OPERATORS *
35 ****
36 0.50 : fracGene (fraction of generation produced by heredity)
37 0.20 : fracRand (fraction of generation produced randomly)
38 0.05 : fracPerm (fraction of the generation produced by permutations)
39 0.25 : fracAtomsMut (fraction of the generation produced by softmutation)
40 ****
41 *          CONSTRAINTS *
42 ****
43 % IonDistances
44 0.6 0.6 0.6
45 0.0 0.6 0.6
46 0.0 0.0 0.6
47 % EndDistances
```

► variable **numGenerations**  
Meaning: Maximum number of generations allowed for the simulation. The simulation can terminate earlier, when the same best structure remains unchanged for **stopCrit** generations.  
Default: 100

► variable **stopCrit**  
Meaning: The simulation is stopped if the best structure did not change for **stopCrit** generations, or when **numGenerations** have expired — whichever happens first.  
Default: total number of atoms for fixed-composition runs, maximum number of atoms maxAt for variable-composition runs.

► variable **bestFrac**  
Meaning: Fraction of the current generation that shall be used to produce the next generation.  
Default: 0.7

► variable **IonDistances**  
Meaning: Sets the minimum inter-atomic distance matrix between different atom types.  
Default: half of the covalent radii sum (in Å) for a corresponding atom pair.



# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## INPUT.txt:

```
34 ****
35 *      VARIATION OPERATORS      *
36 ****
37 0.50 : fracGene (fraction of generation produced by heredity)
38 0.20 : fracRand (fraction of generation produced randomly from space groups specified by user)
39 0.05 : fracPerm (fraction of the generation produced by permutations)
40 0.25 : fracAtomsMut (fraction of the generation produced by softmutation)
```

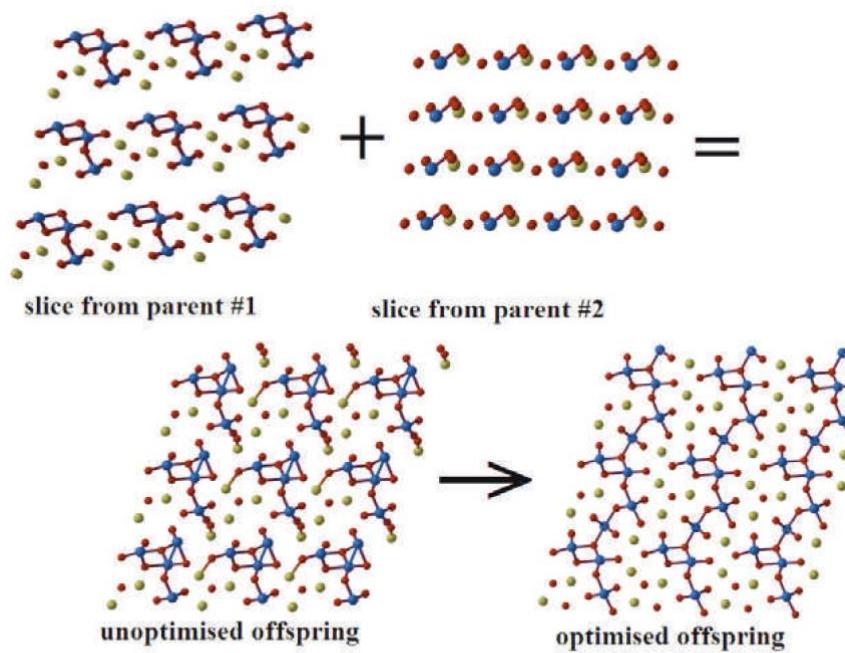


Figure 1. Heredity operator: slices of two parent structures, and the offspring structure before and after local optimization.

Picture from “Oganov A.R., Ma Y., Lyakhov A.O., Valle M., Gatti C. (2010). Evolutionary crystal structure prediction as a method for the discovery of minerals and materials. *Rev. Mineral. Geochem.* **71**, 271-298”

# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

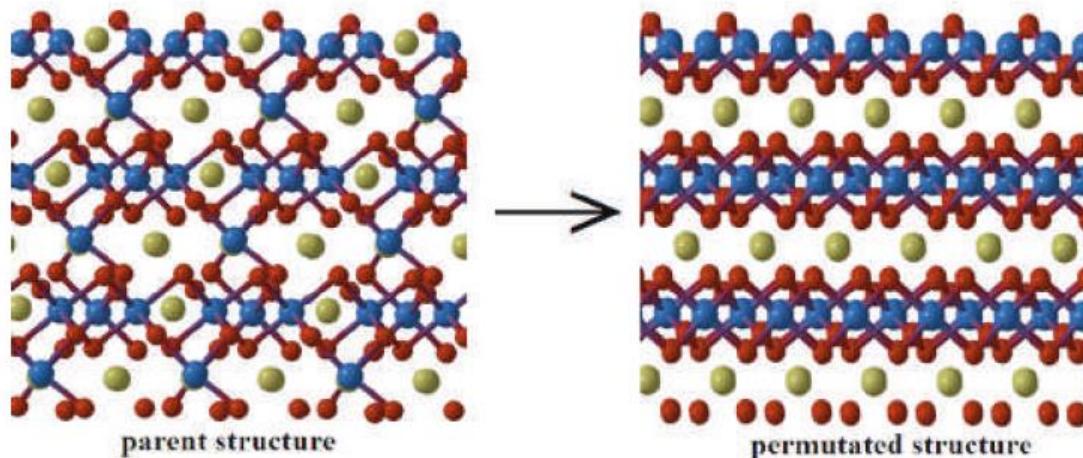
## INPUT.txt:

```
34 ****
35 *      VARIATION OPERATORS      *
36 ****
37 0.50 : fracGene (fraction of generation produced by heredity)
38 0.20 : fracRand (fraction of generation produced randomly from space groups specified by user)
39 0.05 : fracPerm (fraction of the generation produced by permutations)
40 0.25 : fracAtomsMut (fraction of the generation produced by softmutation)
```

▷ *variable fracPerm*

*Meaning:* Percentage of structures obtained by permutation; 0.1 means 10%, etc.

*Default:* 0.1 if there is more than one type of atom/molecule; 0 otherwise.



Pictures from “Oganov A.R., Ma Y., Lyakhov A.O., Valle M., Gatti C. (2010). Evolutionary crystal structure prediction as a method for the discovery of minerals and materials. *Rev. Mineral. Geochem.* **71**, 271-298”

# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

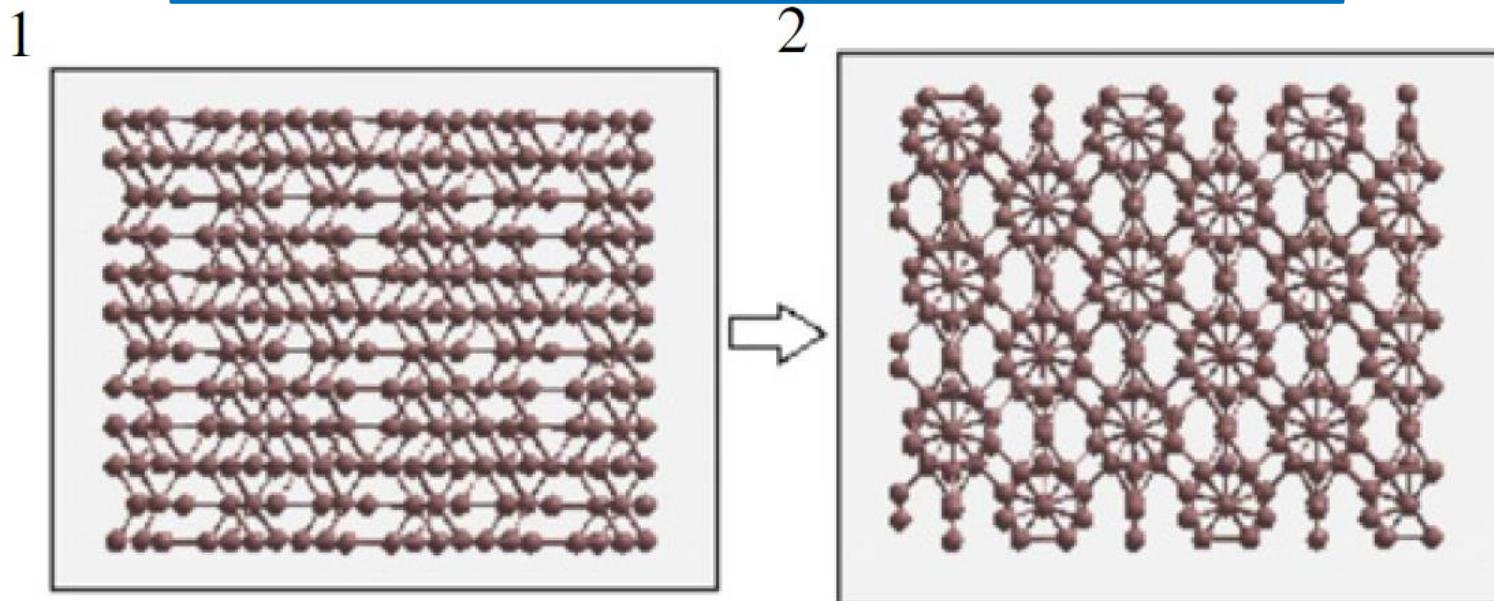
## INPUT.txt:

```
34 ****
35 *      VARIATION OPERATORS      *
36 ****
37 0.50 : fracGene (fraction of generation produced by heredity)
38 0.20 : fracRand (fraction of generation produced randomly from space groups specified by user)
39 0.05 : fracPerm (fraction of the generation produced by permutations)
40 0.25 : fracAtomsMut (fraction of the generation produced by softmutation)
```

▷ variable **fracAtomsMut**

*Meaning:* Specifies the percentage of structures obtained by softmutation or coormutation.

*Default:* 0.1



Picture from “Lyakhov A.O., Oganov A.R., Stokes H.T., Zhu Q. (2013). New developments in evolutionary structure prediction algorithm USPEX. *Comp. Phys. Comm.* **184**, 1172-1182”

# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

In results1/OUTPUT.txt you will notice:

▷ variable `fracLatMut`

Meaning: Percentage of structures obtained from lattice mutations; 0.1 means 10%, etc.

Default: 0 for fixed cell prediction; 0.1 otherwise.

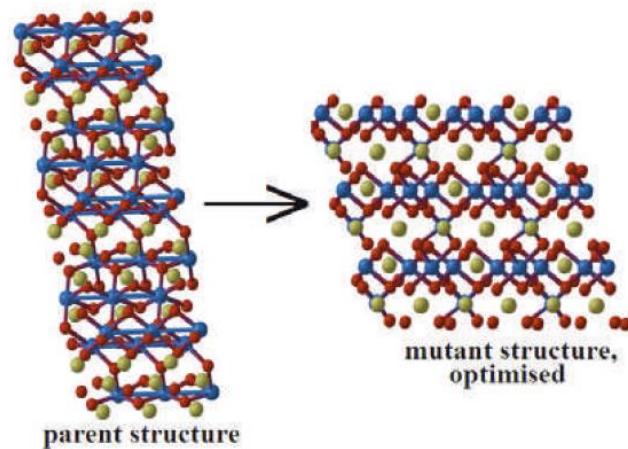
## VARIATION OPERATORS

The fittest 60 percent of the population used to produce next generation

fraction of generation produced by heredity	:	0.50
fraction of generation produced by random	:	0.20
fraction of generation produced by softmutation	:	0.25
fraction of generation produced by permutation	:	0.05
fraction of generation produced by latmutation	:	0.10
fraction of generation produced by rotmutation	:	0.00
fraction of generation produced by transmutation	:	0.00

## Variation operators applied

3 structures produced by heredity
1 structures produced by random
1 structures produced by softmutation
0 structures produced by permutation
0 structures produced by latmutation
1 structures kept as best from the previous generation
0 structures imported from the other USPEX Calculations
0 Seeds structures are added from Seeds/POSCARS



# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## INPUT.txt:

```
51 ****  
52 *      DETAILS OF AB INITIO CALCULATIONS      *  
53 ****  
54 % abinitioCode  
55 3 3 3 ←  
56 % ENDabinit  
57  
58 % KresolStart  
59 0.16 0.14 0.12 ←  
60 % Kresolend  
61  
62 % commandExecutable  
63 gulp < input > output  
64 % EndExecutable  
65  
66 1      : numParallelCalcs (how many parallel calculations shall be performed)  
67 0      : whichCluster (0: no-job-script, 1: local submission, 2: remote submission)  
68 100    : ExternalPressure (GPa)  
69  
70 0      : pickUpYN (if pickUpYN~=0, then a previous calculation will be picked up)  
71 0      : pickUpGen (at which generation shall the previous calculation be picked up)  
72 0      : pickUpFolder (number of the results folder to be used)
```

▷ variable `abinitioCode`

*Meaning:* Defines the code used for every optimization step.

*Default:* 1 for every optimization step (VASP)

*Format:*

```
% abinitioCode  
3 2 2 1 1  
% ENDabinit
```

*Note:* Numbers indicate the code used at each step of structure relaxation: 1 — VASP, 2 — SIESTA, 3 — GULP, 4 — LAMMPS, 5 — Neural Networks code, 6 — DMACRYS, 7 — CP2K, 8 — Quantum Espresso, 9 — ASE, 10 — ATK, 11 — CASTEP.

▷ variable `KresolStart`

*Meaning:* Specifies the reciprocal-space resolution for  $k$ -points generation (units:  $2\pi\text{\AA}^{-1}$ ).

*Default:* from 0.2 to 0.08 linearly



# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

Run the example:

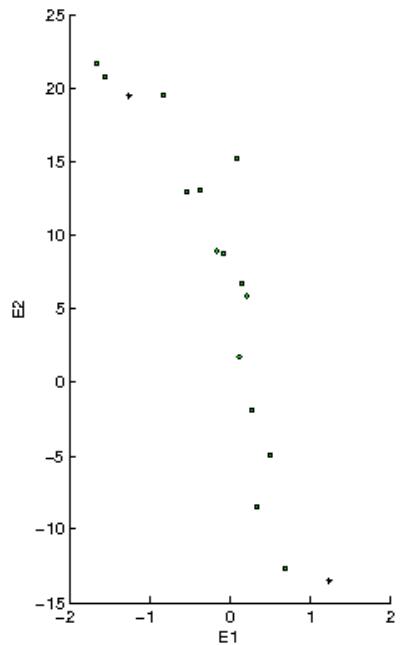
- cp -r ~/examples/example1\_GULP\_300/ .
- cd example1\_GULP\_300/
- bsub < USPEX\_GULP\_300.job

Monitor **log** file, check if **USPEX\_IS\_DONE** file appeared. Then cd to **results1/** to view results.

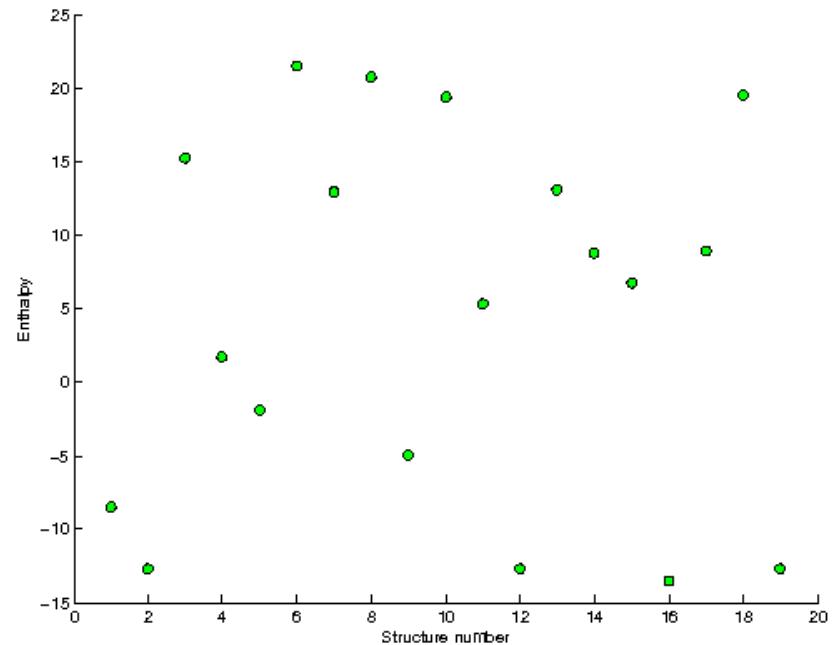
# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## Output files:

E\_series.tif



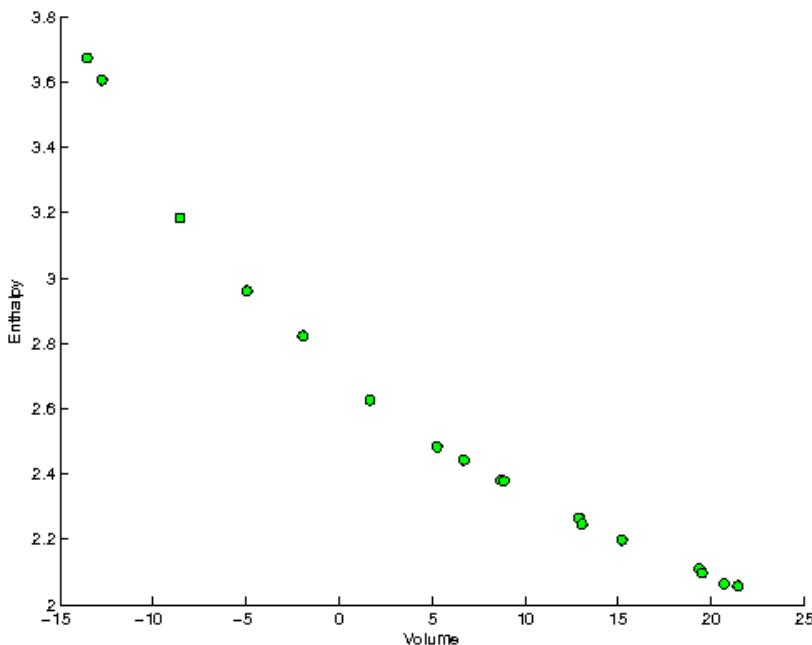
Energy\_vs\_N.tif



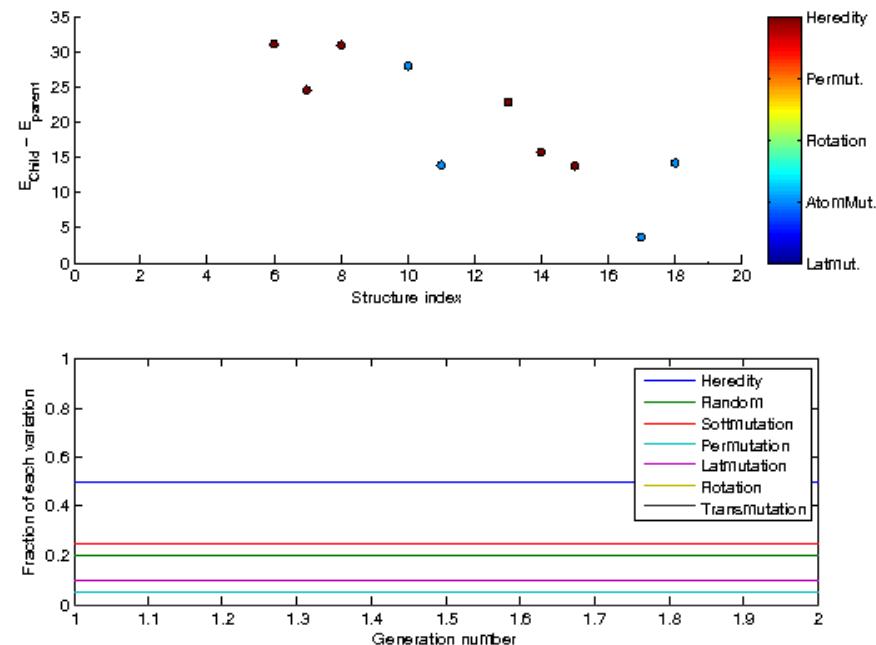
# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## Output files:

Energy\_vs\_Volume.tif



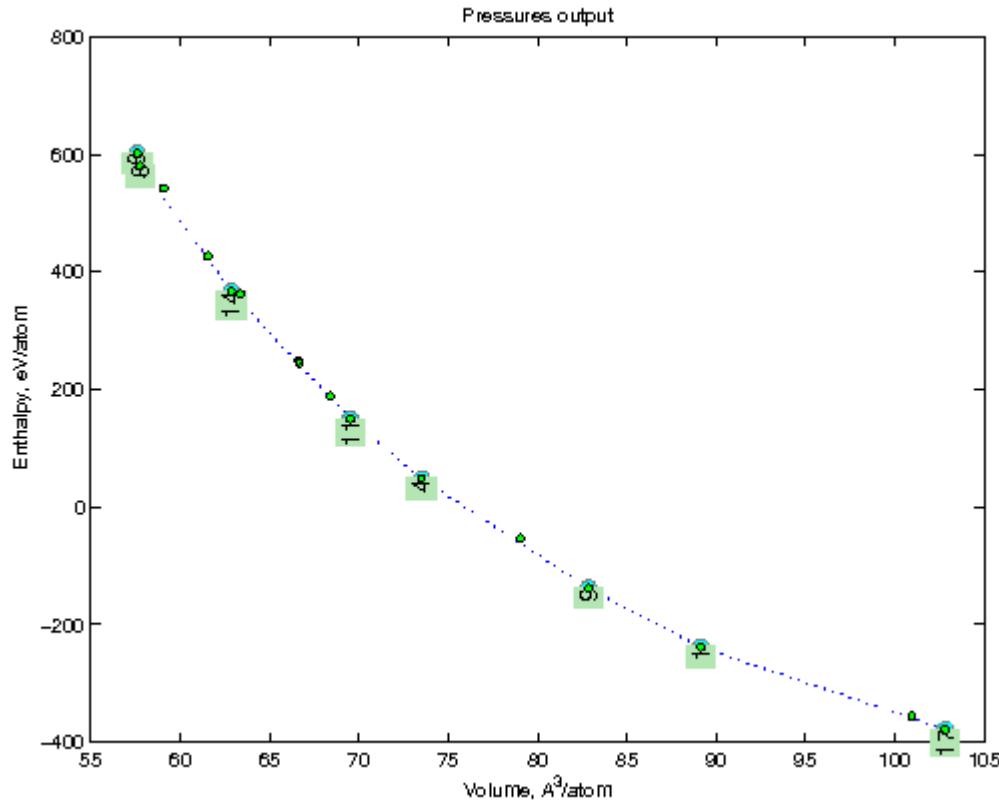
Variation-Operators.tif



# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## Output files:

Pressures\_output.png



# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## Output files:

**log** – contains console output of USPEX

**results1/generation\*/ folders** – contain results in \*.mat files for the generations

**results1/OUTPUT.txt** – contains detailed information on each generation

**results1/Parameters.txt = INPUT.txt**

**Individuals** – summary file where you can find main results:

Gen	ID	Origin	Composition	Enthalpy (eV)	Volume (Å <sup>3</sup> )	Density (g/cm <sup>3</sup> )	Fitness	KPOINTS	SYMM	Q_entr	A_order	S_order
1	1	Random	[ 4 8 16 ]	-238.872	89.138	10.601	-238.872	[ 1 1 1 ]	1	0.207	0.887	1.782
1	2	Random	[ 4 8 16 ]	-356.267	100.939	9.362	-356.267	[ 1 1 1 ]	1	0.244	0.847	1.547
1	3	Random	[ 4 8 16 ]	427.272	61.546	15.354	427.272	[ 1 1 1 ]	8	0.227	0.868	1.645
1	4	Random	[ 4 8 16 ]	47.898	73.509	12.855	47.898	[ 1 1 1 ]	1	0.228	0.831	1.590
1	5	Random	[ 4 8 16 ]	-53.740	79.057	11.953	-53.740	[ 1 1 1 ]	1	0.221	0.807	1.577
2	6	Heredity	[ 4 8 16 ]	601.445	57.609	16.403	601.445	[ 1 1 1 ]	1	0.226	0.774	1.504
2	7	Heredity	[ 4 8 16 ]	362.394	63.374	14.911	362.394	[ 1 1 1 ]	1	0.226	0.801	1.541
2	8	Heredity	[ 4 8 16 ]	580.464	57.730	16.368	580.464	[ 1 1 1 ]	1	0.207	0.812	1.642

**BESTIndividuals** – gives this information for the best structures from each generation

**gatheredPOSCARS** – concatenated POSCAR files

**gatheredPOSCARS\_order** – same as gatheredPOSCARS, but with order parameter

**BESTgatheredPOSCARS** – the same data for the best structure in each generation

**BESTgatheredPOSCARS\_order** – same as BESTgatheredPOSCARS, but with order parameter

# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

## Output files:

**goodStructures** – like Individuals, but ordered by descending of stability

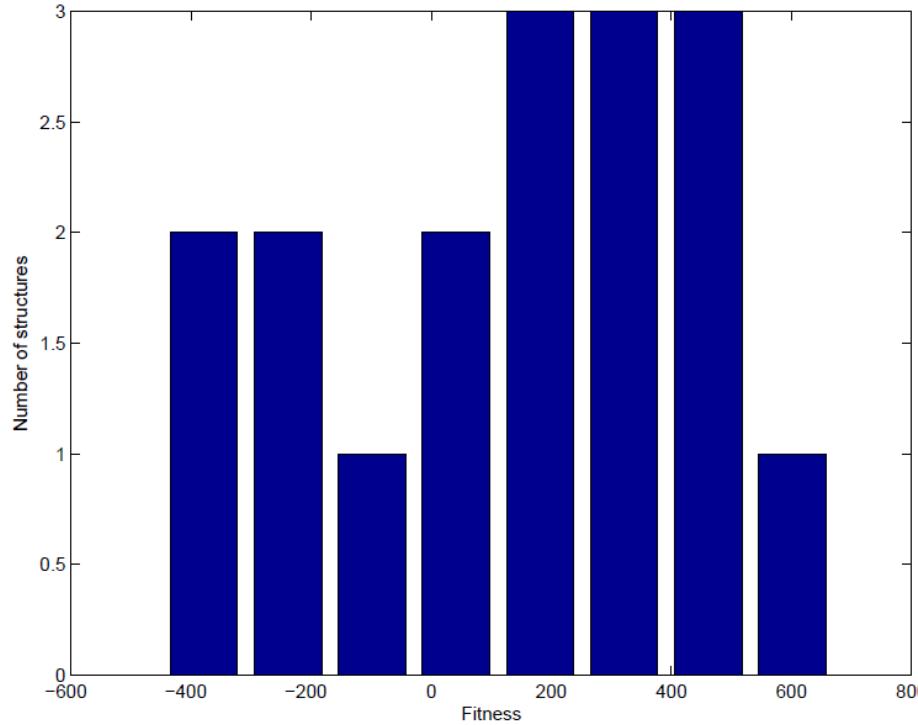
**goodStructures\_POSCARs** – concatenated POSCAR files ordered by descending of stability

**enthalpies\_complete.dat** – gives the enthalpies for all structures in each stage of relaxation

**non\_optimized\_structures** – gives all structures produced before relaxation

**origin** – shows which structures originated from which parents and through which variation operators

**fitnessStatistics.pdf:**



# Bulk Mg<sub>4</sub>Al<sub>8</sub>O<sub>16</sub> (example1\_GULP\_300)

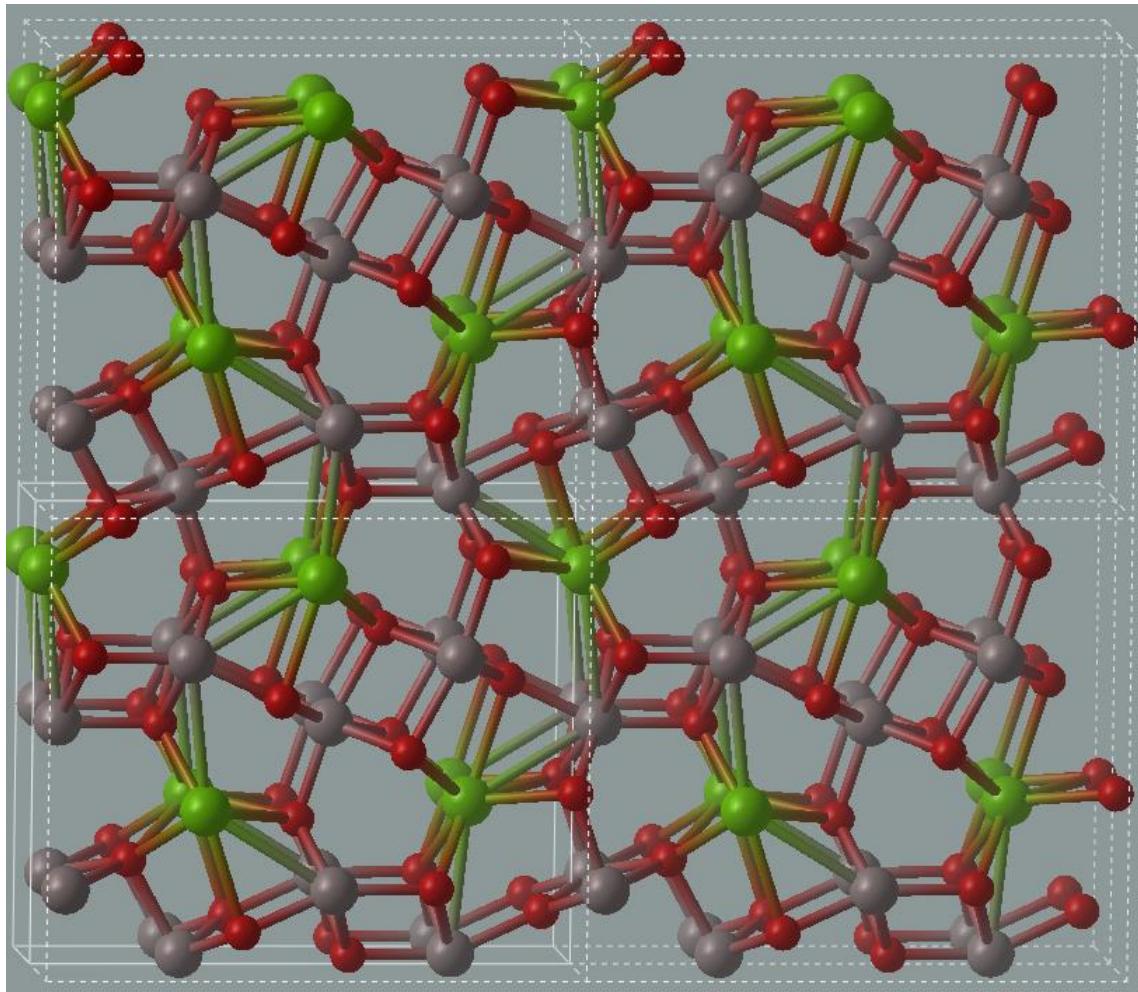
View gatheredPOSCARS:

- cd results1/
- python ~/examples/split\_POSCAR.py

This will produce **POSCAR\_files/** folder with splitted POSCAR files, which you can download on your computer and visualize using VESTA, VMD, etc.

# Bulk $\text{Mg}_4\text{Al}_8\text{O}_{16}$ (example1\_GULP\_300)

View gatheredPOSCARS (from reference\_long/):



# Examples for today:

- Bulk  $\text{Mg}_4\text{Al}_8\text{O}_{16}$  (fixed composition, calculationType=300, GULP)
- Bulk Si-C (variable composition, calculationType=301, VASP)
- $\text{O}_2$  @ Si (100) (fixed composition, calculationType=200, VASP)
- $\text{C}_{13}$  nanoparticles (fixed composition, calculationType=000, VASP)

# Bulk Si-C (example2\_VASP\_301)

## INPUT.txt:

```
1 % PARAMETERS EVOLUTIONARY ALGORITHM
2
3 USPEX : calculationMethod (USPEX, VCNEB, META)
4 301 : calculationType ← dimension: 0-3; molecule: 0/1; va
5 1 : optType (1=enthalpy, 2=volume, 3=hardness, 4=stra
6
7 % symmetries
8 2-230
9 % endSymmetries
10
11 % atomType
12 Si C
13 % EndAtomType
14
15 % numSpecies
16 1 0
17 0 1
18 % EndNumSpecies
19
20 % valences
21 4 4
22 % endValences
```

### ▷ variable `calculationType`

*Meaning:* Specifies type of calculation, *i.e.*, whether the structure of a bulk crystal, nanoparticle, or surface is to be predicted. This variable consists of three indices: *dimensionality*, *molecularity*, and *compositional variability*:

- dimensionality:

- “3” — bulk crystals
- “2” — surfaces, “-2” — 2D-crystals
- “1” — polymers
- “0” — nanoparticles

- molecularity:

- “0” — non-molecular
- “1” — molecular calculations

- variability of chemical composition in the calculation:

- “0” — fixed composition
- “1” — variable composition

*Default:* 300

*Format:*

**Notes:** For variable-composition calculations, you have to specify the compositional building blocks as follows:

```
% numSpecies
2 0 3
0 1 1
% EndNumSpecies
```

This means that the first building block has formula  $A_2C_3$  and the second building block has formula  $BC$ , where A, B and C are described in the block `atomType`. All structures will then have the formula  $xA_2C_3 + yBC$  with  $x, y = (0,1,2,\dots)$  — or  $A_{2x}B_yC_{3x+y}$ . If you want to do prediction of all possible compositions in the A-B-C system, you should specify:

```
% numSpecies
1 0 0
0 1 0
0 0 1
% EndNumSpecies
```

molecular crystal is to be performed, ... with molecular geometries for all in the newly generated structures as 201, -200 (and not yet released: 110,



# Bulk Si-C (example2\_VASP\_301)

## INPUT.txt:

```
29 2      : minAt ←
30 4      : maxAt ←
31
32 0.6    : bestFrac
33
34 0.40   : fracGene (fraction of generation produced by heredity)
35 0.20   : fracRand (fraction of generation produced randomly from space groups)
36 0.20   : fracAtomsMut (fraction of the generation produced by softmutation)
37 0.20   : fracTrans
38
39 2.0    : minVectorLength ( minimal length of any lattice vector) → Only for variable-composition calculations!
40
41 % IonDistances
42 Si  C
43 C  O.0 0.8
44 % EndDistances
45
46
47 abinitioCode (which code from CommandExecutable shall be used for calculation? )
48 1 1 ←
49 ENDabinit
50
51 % KresolStart
52 0.15 0.12
53 % Kresolend
54
55 1      : numParallelCalcs (how many parallel calculations shall be performed)
56 0      : whichCluster (0: no-job-script, 1: local submission, 2: remote submission)
57 0.010  : toleranceFing (tolerance for identical structures)
58
59 0      : pickUpYN (if pickUpYN~=0 , then a previous calculation will be picked
60 0      : pickUpGen (at which generation shall the previous calculation be picked
61 0      : pickUpFolder (number of the results folder to be used. If = 0 , then
62
63 % commandExecutable
64 vasp > output
65 % EndExecutable
```

▷ variable **minAt**  
Meaning: Minimum number of atoms in the unit cell for the first generation.  
Default: No default

▷ variable **maxAt**  
Meaning: Maximum number of atoms in the unit cell for the first generation.  
Default: No default

▷ variable **minVectorLength**  
Meaning: Sets the minimum length of a cell parameter of a newly generated structure.  
Default: 1.8 × covalent diameter of the largest atom.

1 for VASP



# Bulk Si-C (example2\_VASP\_301)

Run the example:

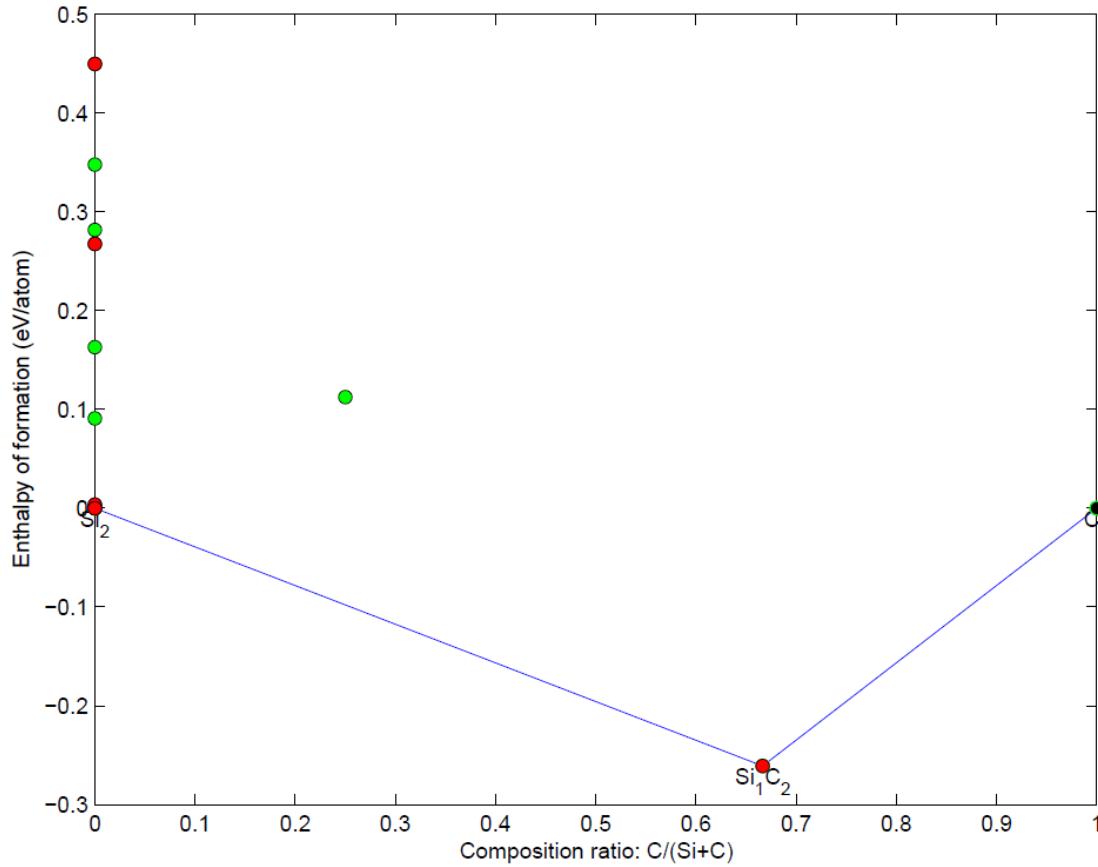
- cd ~/workshop/<your directory>
- cp -r ~/examples/example2\_VASP\_301/ .
- cd example2\_VASP\_301/
- bsub < USPEX\_VASP\_301.job

Monitor **log** file, check if **USPEX\_IS\_DONE** file appeared. Then cd to **results1/** to view results.

# Bulk Si-C (example2\_VASP\_301)

## Output files:

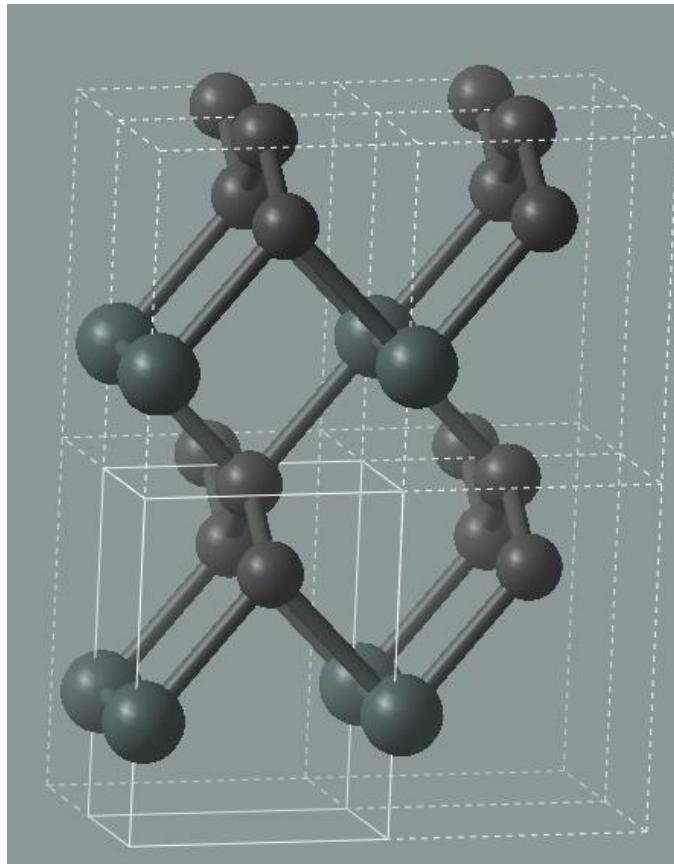
extendedConvexHull.pdf



# Bulk Si-C (example2\_VASP\_301)

## Output files:

**convex\_hull** – gives all thermodynamically stable compositions, and their enthalpies (per atom)  
**extended\_convex\_hull, extended\_convex\_hull\_POSCARs** – all unique low-energy compositions and structures



Your structures  
may be different!

# Examples for today:

- Bulk  $\text{Mg}_4\text{Al}_8\text{O}_{16}$  (fixed composition, calculationType=300, GULP)
- Bulk Si-C (variable composition, calculationType=301, VASP)
- $\text{O}_2$  @ Si (100) (fixed composition, calculationType=200, VASP)
- $\text{C}_{13}$  nanoparticles (fixed composition, calculationType=000, VASP)

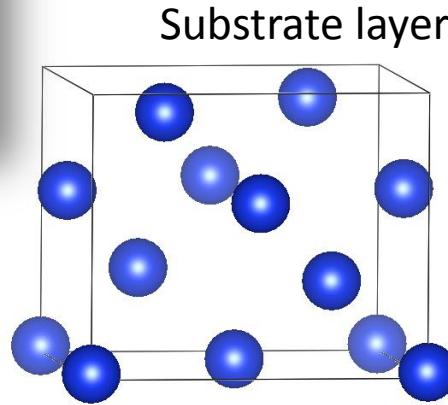
# O<sub>2</sub> @ Si (100) (example3\_VASP\_200)

## INPUT.txt:

```
*****  
*      TYPE OF RUN AND SYSTEM      *  
*****  
*****  
5 USPEX  : calculationMethod (USPEX, VCNEB, META)  
6 200   : calculationType (dimension: 0-3; molecule: 0/1; varcomp: 0/1)  
7 1    : optType (1=enthalpy, 2=volume, 3=hardness, 4=struc_order, 5=aver_dist)  
8 2.0  : thicknessS (thickness of surface region, 2 Å by default)  
9 3.0  : thicknessB (thickness of buffer region in substrate, 3 Å by default)  
10 1   : reconstruct (only used in surface)  
  
11  
12 # vacuumSize  
13 5  
14 # endVacuumSize  
  
15  
16 # atomType  
17 Si O  
18 # EndAtomType  
19  
20 # numSpecies  
21 0 2  
22 # EndNumSpecies  
23  
24 # symmetries  
25 2-17  
26 # endSymmetries  
27  
28  
29 *****  
30 *          POPULATION          *  
31 *****  
32 5   : populationSize (how many individuals per  
33 5   : initialPopSize  
34 4   : numGenerations (how many generations shall  
35 4   : stopCrit  
36 1   : AutoFrac
```

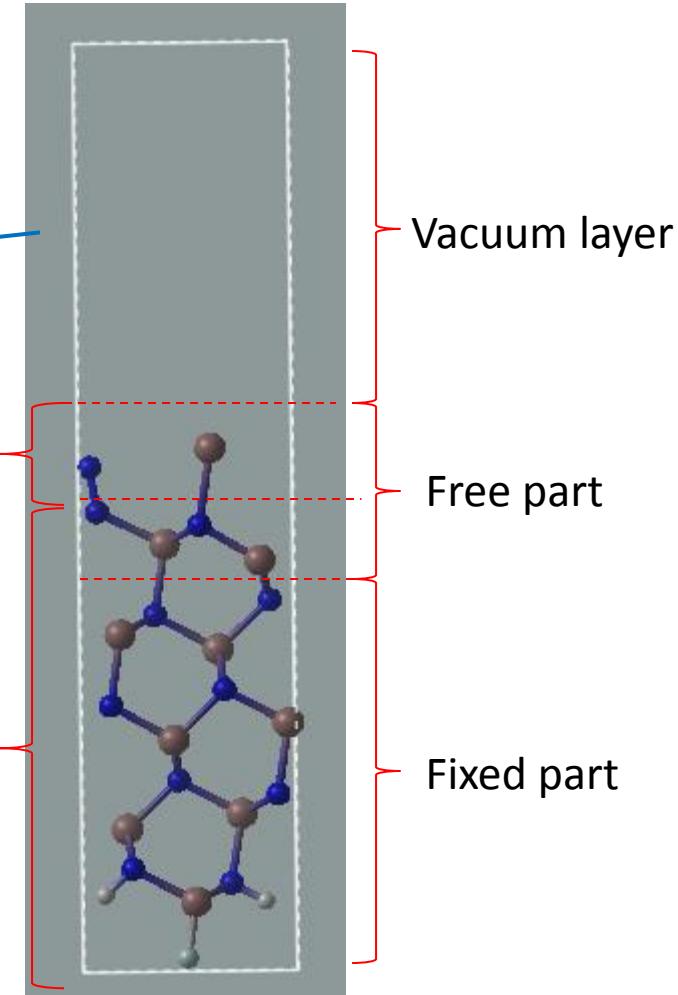
11 Appendix: List of plane groups

Number	Group
1	Group p1
2	Group p2
3	Group pm
4	Group pg
5	Group cm
6	Group cmm
7	Group pmm
8	Group pmg
9	Group pgg
10	Group cmm
11	Group p4
12	Group p4m
13	Group p4g
14	Group p3
15	Group p31
16	Group p31m
17	Group p6
	Group p6m



Add layer

Substrate layer



# O<sub>2</sub> @ Si (100) (example3\_VASP\_200)

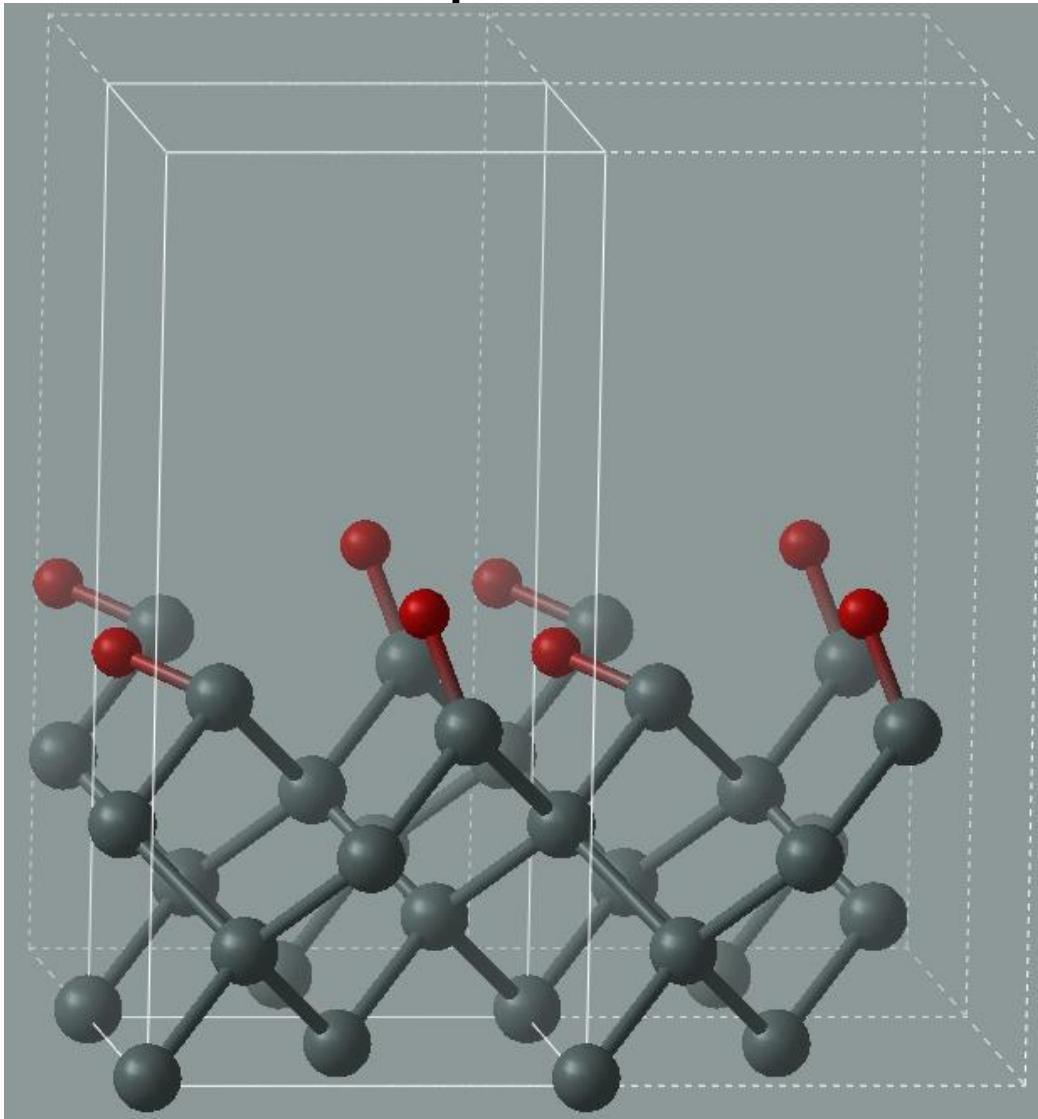
Run the example:

- cd ~/workshop/<your directory>
- cp -r ~/examples/example3\_VASP\_200/ .
- cd example3\_VASP\_200/
- nohup ./uspex\_VASP\_200.sh > out 2>&1 &

Monitor **log** file, check if **USPEX\_IS\_DONE** file appeared. Then cd to **results1/** to view results.

# O<sub>2</sub> @ Si (100) (example3\_VASP\_200)

Output files:



Your structures  
may be different!



# Examples for today:

- Bulk  $\text{Mg}_4\text{Al}_8\text{O}_{16}$  (fixed composition, calculationType=300, GULP)
- Bulk Si-C (variable composition, calculationType=301, VASP)
- $\text{O}_2$  @ Si (100) (fixed composition, calculationType=200, VASP)
- **C<sub>13</sub> nanoparticles (fixed composition, calculationType=000, VASP)**

# C<sub>13</sub> nanoparticles (example4\_VASP\_000)

## INPUT.txt:

```
1 PARAMETERS EVOLUTIONARY ALGORITHM
2 ****
3 ****
4 *      TYPE OF RUN AND SYSTEM      *
5 ****
6 ****
7 USPEX : calculationMethod (USPEX, VCNEB, META)
8 000   : calculationType (dimension: 0-3; molecule: 0/1; varcomp: 0/1)
9 1     : optType (optimise by: 1=enthalpy, 2=volume, 3=hardness, 4=struc_order, 5=aver_dist, 6=mag_moment)
10
11 % vacuumSize
12 5
13 % endVacuumSize
14
15 % What symmetry(s) have to be satisfied by the randomly created structures
16 % symmetries
17 E C2 D2 C4 C3 C6 T S2 Ch1 Cv2 S4 S6 Ch3 Th Ch2 Dh2 Ch4 D3 Ch6 O D4 Cv3 D6 Td Cv4 Dd3 Cv6 Oh Dd2 Dh3 Dh4 Dh6 Oh C5 S5 S10 Cv5 Ch5 D5 Dd5 Dh5 I Ih
18 % endSymmetries
19
20 ****
21 % Here come the atomic numbers of the atoms involved
22 % atomType
23 C
24 % EndAtomType
25
26 % numbers of species (ions/molecules/blocks) of each type
27 % numSpecies
28 13
29 % EndNumSpecies
30
31
32 ****
33 *      POPULATION      *
34 ****
35 5   : populationSize (how many individuals per generation)
36 5   : initialPopSize
37 4   : numGenerations (how many generations shall be calculated)
38 4   : stopCrit
39
40
41
42
43
44
45
46
47
48
49
50
51
52
53
54
55
56
57
58
59
60
61
62
63
64 ****
65 *          CELL          *
66 ****
67 % The following is what you know about the lattice. If you know the lattice
68 % vectors, type them in as 3x3 matrix. If not, type the estimated volume.
69 % For variable composition - type the estimated atomic volume for each element.
70 % Latticevalues (this word MUST stay here, type values below)
71 200
72 % Endvalues (this word MUST stay here)
```



# $C_{13}$ nanoparticles (example4\_VASP\_000)

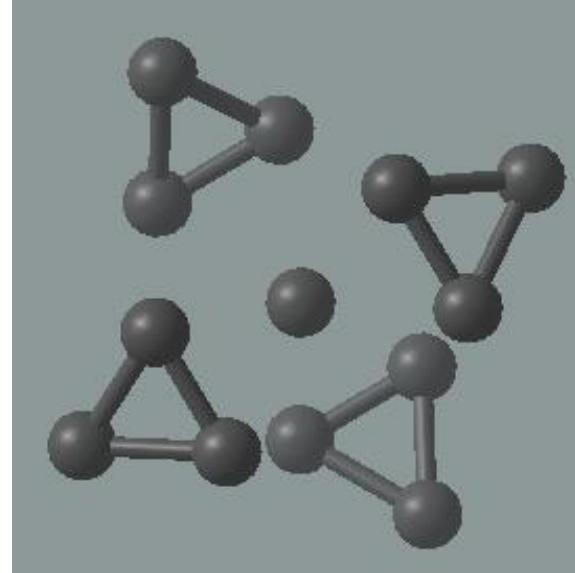
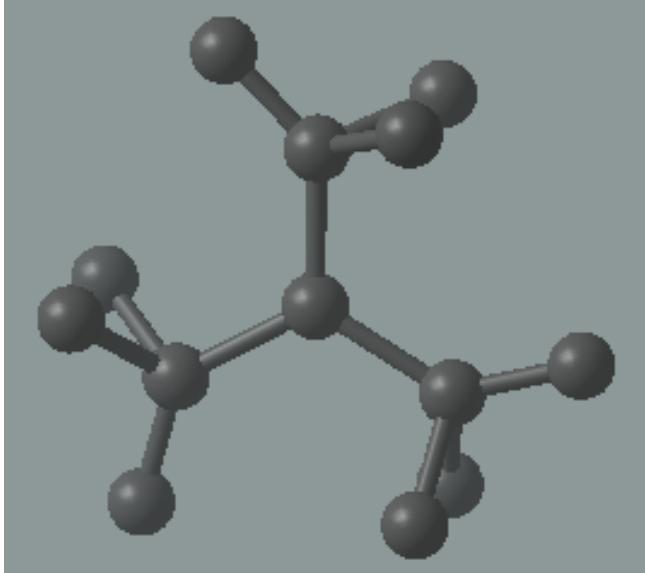
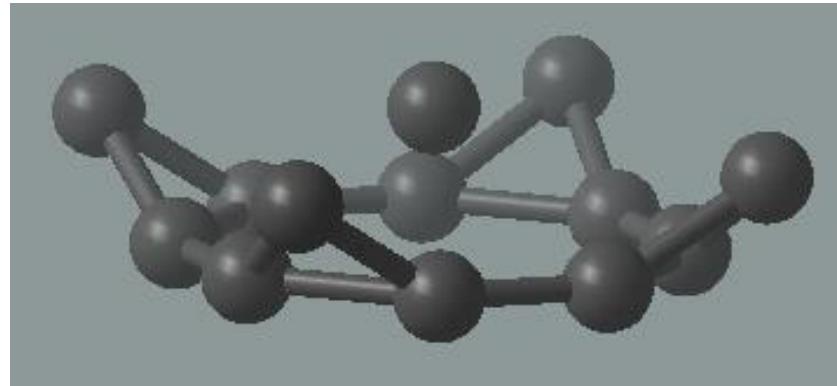
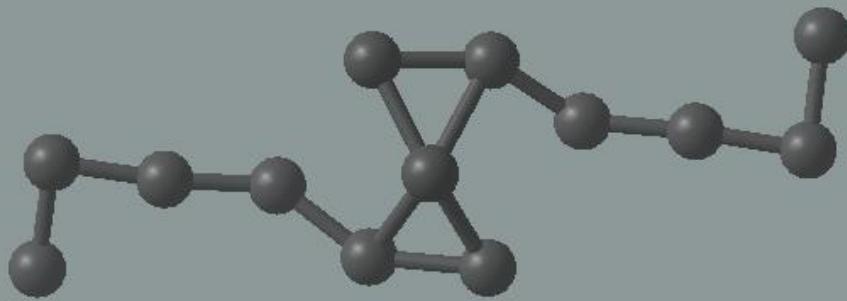
Run the example:

- `cd ~/workshop/<your directory>`
- `cp -r ~/examples/example4_VASP_000/ .`
- `cd example4_VASP_000/`
- `nohup ./uspex_VASP_000.sh > out 2>&1 &`

Monitor **log** file, check if **USPEX\_IS\_DONE** file appeared. Then cd to **results1/** to view results.

# $C_{13}$ nanoparticles (example4\_VASP\_000)

## Results:



**Dank u wel**  
 Pidamayado  
 Dankon

**Shukuria**  
 Wabueja  
 Medawage  
 Merai unachtesh  
**Tingki**  
 Komapsunnida

**Tashakkur**  
 Palidies  
 Haliar  
 Maake Denkaaja  
 Atu gozaimashita  
 Tesekkurer

**Mahalo**  
 Tusen takk  
 Terima kasih

**bolzin**  
 Isi sance  
 bolzini

**Mehrbanı**  
 Nisachalya  
 Ekhmet

**Arigato**  
 Dankscheen

**You**  
 Juspaxar  
 Makatal  
 Balika Yuspagaritam  
 Minnicher Atta  
 Gajitho  
 Yaqhanyelay  
 Efcharisto  
 suksama  
 Shukria

**Merci**  
 Doh je  
 Moiseka skoje  
 Trigipach  
 suksama  
 Shukria

**Spasibo**  
 Efcharisto  
 Salamat

**Grazie mille**  
 Cám ơn  
 Dhanyawaad  
 Köszi

**Biiyan**  
 Toda raba  
 Kamsahamnida

